



SEQUENCE LISTING

<110> DANA-FARBER CANCER INSTITUTE, INC.
KOLODNER, Richard
WINAND, Nena

<120> A METHOD OF DETECTION OF ALTERATIONS IN MSH5

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<141> 1999-12-22

<150> 60/051,686
<151> 1997-07-03

<150> PCT/US98/13850
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<170> PatentIn Ver. 2.1

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Gln Ser Val Val Thr Ser Ala Lys Gln Asp Glu Ala Met Thr Arg Phe
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Leu Gly Lys Leu Ala Ser Glu Glu His Arg Glu Pro Lys Gly Pro Glu
115 120 125

Ile Ile Leu Leu Pro Ser Val Asp Phe Gly Pro Glu Ile Ser Lys Gln
130 135 140

Arg Leu Leu Ser Gly Asn Tyr Ser Phe Ile Ser Asp Ser Met Thr Ala
145 150 155 160

Thr Glu Lys Ile Leu Phe Leu Ser Ser Ile Ile Pro Phe Asp Cys Val
165 170 175

Leu Thr Val Arg Ala Leu Gly Gly Leu Leu Lys Phe Leu Ser Arg Arg
180 185 190

Arg Ile Gly Val Glu Leu Glu Asp Tyr Asp Val Gly Val Pro Ile Leu
195 200 205

Gly Phe Lys Lys Phe Val Leu Thr His Leu Val Ser Ile Asp Gln Asp
210 215 220

Thr Tyr Ser Val Leu Gln Ile Phe Lys Ser Glu Ser His Pro Ser Val
225 230 235 240

Tyr Lys Val Ala Ser Gly Leu Lys Glu Gly Leu Ser Leu Phe Gly Ile
245 250 255

Leu Asn Arg Cys Arg Cys Lys Trp Gly Gln Lys Leu Leu Arg Leu Trp
260 265 270

Phe Thr Arg Pro Thr Arg Glu Leu Arg Glu Leu Asn Ser Arg Leu Asp
275 280 285

Val Ile Gln Phe Phe Leu Met Pro Gln Asn Leu Asp Met Ala Gln Met
290 295 300

Leu His Arg Leu Leu Ser His Ile Lys Asn Val Pro Leu Ile Leu Lys
305 310 315 320

Arg Met Lys Leu Ser His Thr Lys Val Ser Asp Trp Gln Val Leu Tyr
325 330 335

Lys Thr Val Tyr Ser Ala Leu Gly Leu Arg Asp Ala Cys Arg Ser Leu
340 345 350

Pro Gln Ser Ile Gln Leu Phe Gln Asp Ile Ala Gln Glu Phe Ser Asp
355 360 365

Asp Leu His His Ile Ala Ser Leu Ile Gly Lys Val Val Asp Phe Glu
370 375 380

Glu Ser Leu Ala Glu Asn Arg Phe Thr Val Leu Pro Asn Ile Asp Pro
385 390 395 400

Asp Ile Asp Ala Lys Lys Arg Arg Leu Ile Gly Leu Pro Ser Phe Leu
405 410 415

Thr Glu Val Ala Gln Lys Glu Leu Glu Asn Leu Asp Ser Arg Ile Pro
420 425 430

Ser Cys Ser Val Ile Tyr Ile Pro Leu Ile Gly Phe Leu Leu Ser Ile
435 440 445

Pro Arg Leu Pro Phe Met Val Glu Ala Ser Asp Phe Glu Ile Glu Gly
450 455 460

Leu Asp Phe Met Phe Leu Ser Glu Asp Lys Leu His Tyr Arg Ser Ala
465 470 475 480

Arg Thr Lys Glu Leu Asp Thr Leu Leu Gly Asp Leu His Cys Glu Ile
485 490 495

Arg Asp Gln Glu Thr Leu Leu Met Tyr Gln Leu Gln Cys Gln Val Leu
500 505 510

Ala Arg Ala Ser Val Leu Thr Arg Val Leu Asp Leu Ala Ser Arg Leu
515 520 525

Asp Val Leu Leu Ala Leu Ala Ser Ala Ala Arg Asp Tyr Gly Tyr Ser
530 535 540

Arg Pro His Tyr Ser Pro Cys Ile His Gly Val Arg Ile Arg Asn Gly
545 550 555 560

Arg His Pro Leu Met Glu Leu Cys Ala Arg Thr Phe Val Pro Asn Ser
565 570 575

Thr Asp Cys Gly Gly Asp Gln Gly Arg Val Lys Val Ile Thr Gly Pro
580 585 590

Asn Ser Ser Gly Lys Ser Ile Tyr Leu Lys Gln Val Gly Leu Ile Thr
595 600 605

Phe Met Ala Leu Val Gly Ser Phe Val Pro Ala Glu Glu Ala Glu Ile
610 615 620

Gly Val Ile Asp Ala Ile Phe Thr Arg Ile His Ser Cys Glu Ser Ile
625 630 635 640

Ser Leu Gly Leu Ser Thr Phe Met Ile Asp Leu Asn Gln Val Ala Lys
645 650 655

Ala Val Asn Asn Ala Thr Glu His Ser Leu Val Leu Ile Asp Glu Phe
660 665 670

Gly Lys Gly Thr Asn Ser Val Asp Gly Leu Ala Leu Leu Ala Val
675 680 685

Leu Arg His Trp Leu Ala Leu Gly Pro Ser Cys Pro His Val Phe Val
690 695 700

Ala Thr Asn Phe Leu Ser Leu Val Gln Leu Gln Leu Leu Pro Gln Gly
705 710 715 720

Pro Leu Val Gln Tyr Leu Thr Met Glu Thr Cys Glu Asp Gly Glu Asp
725 730 735

Leu Val Phe Phe Tyr Gln Leu Cys Gln Gly Val Ala Ser Ala Ser His
740 745 750

Ala Ser His Thr Ala Ala Gln Ala Gly Leu Pro Asp Pro Leu Ile Ala
755 760 765

Arg Gly Lys Glu Val Ser Asp Leu Ile Arg Ser Gly Lys Pro Ile Lys
770 775 780

Ala Thr Asn Glu Leu Leu Arg Arg Asn Gln Met Glu Asn Cys Gln Ala
785 790 795 800

Leu Val Asp Lys Phe Leu Lys Leu Asp Leu Glu Asp Pro Thr Leu Asp
805 810 815

Leu Asp Ile Phe Ile Ser Gln Glu Val Leu Pro Ala Ala Pro Thr Ile
820 825 830

Leu

<210> 55
<211> 232
<212> DNA
<213> Human

<400> 55
gtaacctccg cgtagacagaa tgagggtggg gcgcgtggag tttccacaa tctgtacttt 60
agttaatac ccgagaattc acctcctgtg tccacagctc tccacgcccc tcagccctgc 120
cccgcagccc tgtatcagaa gtacttagcg cttgcattc tgcgcgccac cctacccgg 180
cctcctctgt gaatcggtgc ttccgaaccg ccctcacttt ttgcattccgc ag 232

<210> 56
<211> 74
<212> DNA
<213> Human

<220>
<221> intron
<222> (73)..(74)
<223> N = A or T or G or C

<400> 56
gtctctgagg ggagtagaaa cttgaatgga gagttgatgg gaatttaaaa taaaagaggg 60
ttgggagccg ggn 74

<210> 57
<211> 189
<212> DNA
<213> Human

<400> 57
aaaaaaaaac agggttggga agagctggc aagtctctta cttcctgagt ggctgttca 60
cattcactaa atgggggtga tcatgcctat ctcagagatt tgagaaatg attaaattat 120
ataagacatg gtaaacccta cacttatgag tgattctaat agtgattcc tttcttcctt 180
gctggacag 189

<210> 58
<211> 450
<212> DNA
<213> Human

<220>
<221> intron
<222> (449)..(450)

<223> N = A or T or G or C

<400> 58

gtggggatgg aaccatgaat tcctctgctc tctgggattt cagatgtgtt acacacacac 60
acacacacac acacacacac acacacatat ttttttttcc tagacagagt ctgtctgt 120
tacccaggct caagtgcagt ggccaaatct tggctcaactg cagcctccac ctccctgggtt 180
caagcaattc tcctgactca acctcccgag tagctggac tacagggctg tgccaccaca 240
cccagctagt ttttgtgtg tggttttagc acagacgggtt tttcaccatg ttggccagg 300
tggctcaaa ctctgaccc tggatccgc ccaccttggc ctctaaagt gctgggacta 360
caggtgtgag tcaccacgccc cagccatgtt ttacttacat taactcacct cactgtctag 420
catatttgtt gttgtgtaa ggaaataacnn 450

<210> 59

<211> 323

<212> DNA

<213> Human

<400> 59

ggcgacaaat atatatgacg tatttacaat gtttcaggtt cttcagattc agccctgggc 60
aaatcagtca tgcgtgttct ccaggggttt acagcctagt gacaacatcc agaacatccc 120
acttccctct caccatccca ccactctaa ctactttctt aaatctcaac ttctacctgt 180
gttcccactg tgcagagcac tccctactcc tagggaggaa atgttttga gaaggagagg 240
ggtaggaaga ggagggctat gggtttctc ttagtcaaag acaaagatcc tttaactcat 300
ttgatctctg ttctccttcc aag 323

<210> 60

<211> 150

<212> DNA

<213> Human

<400> 60

gtaaggactt ggtaaaggat agagggaaaa tggggaaagga ctaatataatg gaatattcca 60
gggggctaga atttgggtgag agggagtgac agacagaggt agaaggactg agatgtaaag 120
aatgatagcc ttttcttcc tcccccacag 150

<210> 61

<211> 733

<212> DNA

<213> Human

<400> 61

gtatctcctt cctttgttt tgcctaactc cctgttccgg tgccttattt tttcccccaa 60
ctctacccctt atcatcacag atctcccttc tgccttattgt catcctaaac ctttgtctc 120
ctcatgcctt atgacctgtc ccccaagat ctctcctgtc ccctaccctt taataatctg 180
cagcttattt ggaaggctt gcttaagtca tgcgttaggaa tgaggccctc ccctgaggag 240

tggtgacact ttttggacag ggttttattt tggaattct ccccatthaag ttaaagcctt 300
ttatcaccaa accaaaaggc actgcctcag tgacccttat tatgatccat aaggcacttc 360
tataactttc ctaggtttac aataagaaca ggagtgtact atcctaatta gatattaagg 420
cattagtggtt actagttcta ttaataccat tattttgacc aaaatcctca attccagaca 480
gatgtctact ttccctcagcc atttatcttt ctcaggctgt gcttcagac aagtatctt 540
atattatatg tagaataaaaa agagaattag actaagagtc tgaaaatttgc ttcttgctc 600
tagctttcca ttaactgcct gtgtgagctt gggcaagtca aataatctct ctgccttcta 660
ttgtctcatt cttaaaaatgg ggtgaaaaaa ttgagctaca agaccgttcc ctggcttgc 720
ctccctcaaa tag 733

<210> 62
<211> 164
<212> DNA
<213> Human

<400> 62
gtgagattgg tcctggggga taagggctgg gaggcggcac aagtgcattt gctgaattct 60
gggaggtact ggcttagccc tggaaaatag taactttccc ttgtgctctg cagccccag 120
gagatttaag atttaccccg attccactgc tgatcccctc ccag 164

<210> 63
<211> 246
<212> DNA
<213> Human

<400> 63
gtaggtgatt caccccaacc ccaaccaaag taatgtggga ttggggggcc tgaaaagtaa 60
agtgggggtg ggggtggat gtggctgtga cccagtgggt caagggtctt agacaccccg 120
ggagaatcta agggctaattt agactttggg aagaagactg ggacaatatt cagagagggg 180
gacaaaggaa gtggagttgt ggaacgaact cagactgcctt cctgctttt tgggggtgt 240
cctcag 246

<210> 64
<211> 413
<212> DNA
<213> Human

<220>
<221> intron
<222> (412)..(413)
<223> N = A or T or G or C

<400> 64
gtaaagaggt ggaggcatgc tgctgtctt ggggagggag aaggattaag ttatgcctt 60
caataatctt aatggggctc tagttccctt aatccctgggg ctatataatg ctctctcctt 120

gaaggaaagg gaaggggggt tttgagggaa agagaggaag aaaagcataa agataactagc 180
tttctttct atagggagaa actgaggcaa agaaaaagtaa gggacaaacc ttacatcaag 240
atatgatctc ggctgggcgc ggtggctcat gcctgtaatc cccgcgttt gggaggccaa 300
ggcgggtgga tcgcctgagg tcaggagttt gagacctgac caatatggta aaaccccgac 360
tctactaaaa atataaaaaat tagctgggtg tgggtgcgc ctgtaatccc ann 413

<210> 65
<211> 136
<212> DNA
<213> Human

<400> 65
tttttttta aaaaaaaaaa aaaaaagacg tgatctcagg aggatatccc ctgtcccat 60
tccatttatac agtctcaat tcttattccc ctaaaaagtc caagttaccc caaactcctc 120
catttctcct cgacag 136

<210> 66
<211> 356
<212> DNA
<213> Human

<220>
<221> intron
<222> (355)..(356)
<223> N = A or T or G or C

<400> 66
gtaggtgtgc cccatccctc atctcacgta caaagaccta ccagaaaagc aattggctcc 60
aaagatgtgt cccagcctcc ctcccactt cactcccatt gtcagatatac tcttcatgc 120
caatccaaat ttcttaccta ttgtacccc ccgcggggca agcttgagca tcttccata 180
ctttgtggct gtacagtgtg ttgcataatca gccattactt taccattct gtgtccctc 240
cctgggtttg tatgaatgtt tctactagtt gggtagctgt tagggacttt ggagacctt 300
gtgtatagag aagagtttg taactgcata actgcctatt tgattttat agagnn 356

<210> 67
<211> 426
<212> DNA
<213> Human

<400> 67
ccaggagtag agggagagac agaaacagcc aacaatggcc cagaaaatgg atgatataatt 60
agataagggaa agaaaatgagt taccagattt gggagagatg gtttggatgt caaagcaggt 120
gatcggtgac gtcagcgtcc gagggaaagac ggctgccacc ggccggggca gttgagggaa 180
ctaggttagtt aagtgtgtc gggctaaaag tccctagagt gtccatccct ccccccattc 240
catgtgcgtt aatcccagct cattttagggg ccaggcacca actttgggtt ccttggcc 300

ctcccaggcc agttcctca acaaccagca cctctgactg gatgcctcg gtagacaca 360
taaacacatt ccattgcct gtccgtgcct tgtaacaagt tcactccctg ccttatccct 420
cacaag 426

<210> 68
<211> 360
<212> DNA
<213> Human

<220>
<221> intron
<222> (359)..(360)
<223> N = A or T or G or C

<400> 68
gtgagtgggt cccacacata ctacacacta atgcatgaat tccatatgca cactacatac 60
taagcctact aatggcagta tacagattct cacatacacc accccaccta gtagtagtaa 120
agcaactgccc cttactgag cactggctaa ctgcattca tccttataac agctttgtgt 180
agtagctgat atgcatctca tttttgttg tcagcgcagg tacacatata cattgatgat 240
acacagactt gcacacatac agcagcagga aaaaacacaa aatgtaaggc cgggcacagt 300
ggctcacacc tgttatcagc actttggggg gccaacgctg ggtgacccctc catcttggn 360

<210> 69
<211> 447
<212> DNA
<213> Human

<400> 69
cacaggaaga atatgaaaag atgaatgtct gttgctgtta cccagagaca ctttcacagc 60
taaaaagaca tacaactca tactgactca ccgtctctta ctcagcctca gagttagctg 120
cagtgttggc acacaaatac ctcaacacac tgctctcctt ctaaaatatt gacaagctcc 180
gttacttata tacatggaat gacacacggt cttatccgtt gaaactgtga tatgttagaca 240
caattatgtc cacatcttagc aattttcagt agatacatgt aaacacaccc gaatgggttag 300
gacactgcac ttgccactac attcccatag cacatcgtgg atacatattg ccacaatccc 360
cagggactgc aagcacactt tttggcaaac tgagatcaag atgatagatg taactttag 420
taccccccacc caaaccctca cttccag 447

<210> 70
<211> 127
<212> DNA
<213> Human

<400> 70
gtgagcccaag ggtggagggc agggagggtgg ggaaggaggt tgagggctga tactggcag 60
tgggcttctt gagggcatt agagtggagg aagagaaaac agcggctgta accttgcgt 120

actgttag

127

<210> 71
<211> 30
<212> DNA
<213> Human

<220>
<221> intron
<222> (29)..(30)
<223> N = A or T or G or C

<400> 71
gtaaggcatt ccttcttgaa tcccaaaaann

30

<210> 72
<211> 222
<212> DNA
<213> Human

<400> 72
tacaggcatg agccactgtg cctggccagg accatatctt aattgtcttt gtagttcag 60
tgtttggtagt agtgcctctc actgtttctt tttgccttg agatctccc tccttggtag 120
tgtgatcttc cctactggtc ttgttcttc tgagtctgtc cctatcacca cctcaacccg 180
agctggatgt ggcctgtcct ccttttgtg tttctctcac ag 222

<210> 73
<211> 254
<212> DNA
<213> Human

<400> 73
gtgagtagaa ggaaaaaggg agtgcaccca gggaggtcag ggagagagaa tgcaagtgtgc 60
aagatgggaa aacatgaaag atattgaggt caattggata aagaatggaa tggggagg 120
aggcagcaga acttcaggaa agtatctgga gggtagagt taaaggagga ctgcaggag 180
aattggggcc caaggagagc tgaggaacag gacagagggt gccaggtcct aagaaacagt 240
acttatctcc tcag 254

<210> 74
<211> 145
<212> DNA
<213> Human

<400> 74

gtgagtgttg ggtgtggatg ggcctgttag ccctgcgcag ttagggatg ccatccttgg 60
caggtggta ccacagctgg ggatcttcat agcaaccagg gcaggagact cactttgat 120
aaccacctgt ctccaccct cgtag
145

<210> 75
<211> 98
<212> DNA
<213> Human

<220>
<221> intron
<222> (97)..(98)
<223> N = A or T or G or C

<400> 75
gtgagggcag gagagtgggt gtagcctca gatgtctttt gggggagata ttaggcttat 60
gaaagacata ctggtagata agaaaacttg tggggcnn
98

<210> 76
<211> 83
<212> DNA
<213> Human

<400> 76
atcttttaag ctcccttggg atggggaggt tccagtaagt ctccaaacaa gagagttagag 60
tatctcctct ttactctccc cag
83

<210> 77
<211> 247
<212> DNA
<213> Human

<400> 77
gtaagaccct caacctctgt aaggtagtg atgaggaaaa ttagtcagca gctgaggaag 60
agcgttactc tacagcagca ctgccaata tggatctct cctctgttagt tttactctga 120
gctttaccag cactgagaca aaggaaagag aagtcagagt tagggctgg aggtgggtt 180
agaaagatgg ggaaggagag gaggaccaag agatgcaaag tccacagctt tgaaccctg 240
tacccag
247

<210> 78
<211> 273
<212> DNA
<213> Human

<400> 78

gtgaggaaaa gccagagggtt atatgcattg taagatgttt aaaaaaagca gcagccaggg 60
gaaggagggg agtgggcaac ttggggatgc ttccaacagg cccctctct tcctgctctc 120
tgtctcgctc actctgactc tatctttcc tctgaatgtc ttgaggctc agattgtatc 180
tgcaacctgt ttccagatcc ccctagggc ctctgcctct cttcaactt cccctggAAC 240
tgacctccag ctcccttcctt caccactcc cag 273

<210> 79

<211> 114

<212> DNA

<213> Human

<400> 79

gtaagaatag aggccgggtgg aggaatacac atgaggggcc caaaggctac atcttctggg 60
ggttcatcta tcttgatcca caagccatgc gaggtgcctc tccgcccact gcag 114

<210> 80

<211> 473

<212> DNA

<213> Human

<400> 80

gtgaggagaa gccctgcagc ctgggcctct ggctgtctct gcatctactc caccctact 60
tgccagccaa ctcaggctcc tgcagctctt ctcccatttt ctgaccggc tcttcatgaa 120
aggaccatca cccacatccc tggcttcca cctcacatgt tcttattctc cactggagag 180
ccatgctcta atgaaactt ccgtggccca aattcctca cctgcctctg agtaggtaca 240
caccactccc aagtatgtct ctgcccacgt cccgtgcctc ttcaactgatt ctaaatttagc 300
ccacagggct atggtcagga ttggggagg agagacagag tcagtgtgtc tggcacat 360
ttctcctgtt tcaccctgtc catttctctt tgatgtgcca ttcatgcctt gaggctact 420
ttcacctcag cccacggcac caggccccag gccctgtctc cttccctatt cag 473

<210> 81

<211> 348

<212> DNA

<213> Human

<400> 81

gtcaaaggga acaaaggag gtgggattga ggaagggat aatggaaag gaaccctga 60
aaatgctcat aacaggaaag catgcctct gctgcattgc cttaatacta aaagtggga 120
gcactaaagg cagagataag aagaatcaat accataaaca ttcttgaac ctttgcgtca 180
tgtgagtcac tggggcaaa gaggatgaac aaagcgtgca cttcaccatt caagaacttg 240
cagtgcagta gggagggcat gtatacagct ttattcacag gccaactgtg gtcagtgcgt 300
tacgggcttc caataactaac ttccccttgt ccaccttata cccagcag 348

<210> 82
<211> 209
<212> DNA
<213> Human

<400> 82
gtgaggggag aaactgatga ggggagaac taaggagggg aaaatggagg aggatgaagg 60
agcatgacag tgaggctggg cctctgaat ggaatagggc tgtgtggca gaaaagaaaat 120
agaacacgag acagggaaag gcagtgcag tgcaaggggg catatgggt cccatggct 180
ccgaatgcta acctctgccc tctttgcag 209

<210> 83
<211> 202
<212> DNA
<213> Human

<400> 83
gtgaggagac caatcttagct cctcggggac ccccaggctg ggcatttccc agaggtgggg 60
atggctccct ctatcagaac aagggctccc tcagcacaga gaccacatcc ctccctttt 120
ctccctcccc acaggattgg ccaagggttt caggacagga aggaggtgat tcatgataca 180
ctgtctttta ttctctttta ag 202

<210> 84
<211> 155
<212> DNA
<213> Human

<400> 84
gtgatgagat ccaaattgtgc aaccacctcc acatcagagc tcccttcat tcctagtct 60
actgggcctg ggtcttaggtc cacaggattt ctgaccctta tttcccttc tcttccccac 120
tccccttact cctcccacct tcttgcttgt cctag 155

<210> 85
<211> 215
<212> DNA
<213> Human

<400> 85
gtgcgtatat ggccccagtg tctttaccct ctctgcatct tctcctgcaa ctcttctccc 60
ccctccagca cttingccctt cagaaaccca ccatttcttt ctgaaatccc taaatcttca 120
agatcccagg ttttctgtgc cacagccctt cccctctgcc cagggatttg gtgtccatt 180
ctgccataaa tcttgcgatt ttctcttcc tttag 215

<210> 86

<211> 29
<212> DNA
<213> Human

<400> 86
gctgctcagg tatacagtagc cacgctccc

29

<210> 87
<211> 29
<212> DNA
<213> Human

<400> 87
agatccgggg tgaggagccc gtggtagga

29

<210> 88
<211> 29
<212> DNA
<213> Human

<400> 88
gaatggcagg tgagaagggg ccccatgtc

29

<210> 89
<211> 29
<212> DNA
<213> Human

<400> 89
ctcaagcagg tgaggggccc ccaagctgg

29

<210> 90
<211> 29
<212> DNA
<213> Human

<400> 90
accaactcgg tgcgaggaa aatgaagag

29

<210> 91
<211> 29
<212> DNA
<213> Human

<400> 91	
ttcccatccc aaccctccag gctgtggtt	29
<210> 92	
<211> 29	
<212> DNA	
<213> Human	
<400> 92	
ctctctct ctttccag accaggaga	29
<210> 93	
<211> 29	
<212> DNA	
<213> Human	
<400> 93	
tgtctctcta cccaccacag gcattct	29
<210> 94	
<211> 29	
<212> DNA	
<213> Human	
<400> 94	
tctccctgc cttggccag gtggctt	29
<210> 95	
<211> 29	
<212> DNA	
<213> Human	
<400> 95	
tcacctctgc ctttgacag gtggatggc	29
<210> 96	
<211> 79	
<212> DNA	
<213> Human	
<400> 96	
gtatacagta ccacgctccc caagcaaagt caagatgaga gaagacgtga ctgttaacct	60

tcccatccca accctccag

79

<210> 97
<211> 135
<212> DNA
<213> Human

<400> 97
gtgaggagcc cgtggtagga gggggcaggc tgctctaaca gaccctgctc tcatgctggc 60
ccctctgcat ggtcacactg catctgcatt cctgcttcca gatcttcca ggcacctctc 120
tctctcccttc tccag 135

<210> 98
<211> 79
<212> DNA
<213> Human

<400> 98
gtgagaaggg gccccatgtc ctgctgtggg gatcctccct gggtccacaa accatgcagt 60
gtctctctac ccaccacag 79

<210> 99
<211> 389
<212> DNA
<213> Human

<400> 99
gtgaggggcc gccaagctgg gggcccacat ctccatctcc tctggccgcc aggccagatc 60
ctctgcccc ccccacacac acatacagca catgtccttg tcctctgagg gacagtctgt 120
tcttttagat agacctttcc gtggccacaa gtccctggac caacctccaa atagatccat 180
gccgttccct agtatgcctt taccacaaac cttgactctg gagttattg tgaagtcaagg 240
acccaggaaaa ctgtgttcca gggctctgtt cttctgttac actgtgtcct ctcttaatc 300
tgtcggtcat gtcttagtt gagaccatt tttactttgc ccatagtagc gcaacaggcc 360
catgttctgt ctccccgtcc ctggcccaag 389

<210> 100
<211> 180
<212> DNA
<213> Human

<400> 100
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21

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28